SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i)	APPLICANT:	Sherrol H. McDonough
		Daniel L. Kacian
		Nanibhushan Dattagupta
		Diane L. McAllister
		Philip Hammond
		Thomas B. Ryder

(ii) TITLE OF INVENTION: NUCLEIC ACID SEQUENCE AMPLIFICATION

(iii) NUMBER OF SEQUENCES: 23

(iv) CORRESPONDENCE ADDRESS:

(A) (B)	ADDRESSEE: STREET:	Lyon & Lyon 633 West Fifth Street
(C)	CITY: STATE:	Suite 4700 Los Angeles California
(E)	COUNTRY:	U.S.A. 90071-2066

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: IBM B C DOS 5 0

(C) OPERATING SYSTEM: IBM P.C. DOS 5.0 (D) SOFTWARE: Word Perfect 5.1

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 08/480,472 (B) FILING DATE: June 6, 1995 (C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 08/345,861 (B) FILING DATE: November 28, 1994

(A) APPLICATION NUMBER: 07/925,405 (B) FILING DATE: August 4, 1992

(A) APPLICATION NUMBER: 07/855,732 (B) FILING DATE: March 19, 1992

(A) APPLICATION NUMBER: 07/550,837 (B) FILING DATE: July 10, 1990

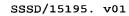


	(A) APPLICATION NUMBER:(B) FILING DATE:	07/379,501 July 11, 1989
(viii)	ATTORNEY/AGENT INFORMATION	ON:
	(A) NAME: (B) REGISTRATION NUMBER (C) REFERENCE/DOCKET NUI	Heber, Sheldon O. : 38,179 MBER: 213/066
(ix)	TELECOMMUNICATION INFORMA	ATION:
	(A) TELEPHONE: (B) TELEFAX: (C) TELEX:	(213) 489-1600 (213) 955-0440 67-3510
(2) INF	ORMATION FOR SEQ ID NO: 1	:
(i)	SEQUENCE CHARACTERISTICS	:
	(C) STRANDEDNESS: sir	base pairs cleic acid ngle near
(ii)	SEQUENCE DESCRIPTION: SEQ	ID NO: 1:
GAAATTAA AAGCT	TA CGACTCACTA TAGGGAGACC AC	CAGCCGTCA CCCCACCAAC 50
(2) INF	ORMATION FOR SEQ ID NO: 2:	
(i)	SEQUENCE CHARACTERISTICS:	
	(B) TYPE: nuc (C) STRANDEDNESS: sir	base pairs cleic acid agle near
(ii)	SEQUENCE DESCRIPTION: SEQ	ID NO: 2:
GGGATAAG	CC TGGGAAACTG GGTCTAATAC C	31
(2) INFO	ORMATION FOR SEQ ID NO: 3:	
(i)	SEQUENCE CHARACTERISTICS:	
	(B) TYPE: nuc (C) STRANDEDNESS: sin	base pairs leic acid gle lear
(ii)	SEQUENCE DESCRIPTION: SEQ	ID NO: 3:

(2) IN	FORMAT	TION FOR SEQ ID	NO: 4:	
(i) SEQ	QUENCE CHARACTER	RISTICS:	
	(B) (C)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	23 base pairs nucleic acid single linear	
(ii)	SEQU	JENCE DESCRIPTIO	ON: SEQ ID NO: 4:	
CCGGATA	GGA CC	CACGGGATG CAT		23
(2) IN	FORMAT	TION FOR SEQ ID	NO: 5:	
(i)	SEÇ	QUENCE CHARACTER	RISTICS:	
	(B) (C)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	20 base pairs nucleic acid single linear	
(ii)	SEQU	JENCE DESCRIPTIO	N: SEQ ID NO: 5:	
CGGTGTG	GA TG	ACCCCGCG		20
(2) IN	FORMAT	ION FOR SEQ ID	NO: 6:	
. (i)	SEQ	UENCE CHARACTER	ISTICS:	
	(C)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:		
(ii)	SEQU	ENCE DESCRIPTIO	N: SEQ ID NO: 6:	
AATTTAAT	'AC GA	CTCACTAT AGGGAG	ACCA GGCCACTTCC GCTAACC	47
(2) INE	ORMAT	ION FOR SEQ ID	NO: 7:	
(i)	SEQ	UENCE CHARACTER	ISTICS:	
	(A) (B)	LENGTH: TYPE:	24 base pairs nucleic acid	

	(C) (D)	STRANDEDNESS: TOPOLOGY:	single linear	
(ii)) SEQU	JENCE DESCRIPTION:	SEQ ID NO: 7:	
CGCGGA	ACAG GO	CTAAACCGC ACGC		24
(2) II	NFORMAT	TION FOR SEQ ID NO	8:	
(:		QUENCE CHARACTERIST		
	(A) (B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	23 base pairs nucleic acid single linear	
(ii)	SEQU	JENCE DESCRIPTION:	SEQ ID NO: 8:	
GGAGGA'	ratg to	CTCAGCGCT ACC		23
(2) TI	VEORMAT	TION FOR SEQ ID NO:	· 9:	
		QUENCE CHARACTERIST		
(-	-			
	(B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	nucleic acid single linear	
(ii)	SEQU	JENCE DESCRIPTION:	SEQ ID NO: 9:	
CGGCTG	AGAG GC	CAGTACAGA AAGTGTCGT	TG GTTAGCGG	38
(2) II	NFORMAT	TION FOR SEQ ID NO:	: 10:	
(:	i) SEÇ	QUENCE CHARACTERIST	TICS:	
	(B)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	36 base pairs nucleic acid single linear	
(ii)	SEQU	JENCE DESCRIPTION:	SEQ ID NO: 10:	
GGGTAA	CCGG GI	AGGGGTTG TGTGTGCGC	GG GTTGTG	36
(2) II	NFORMAT	TION FOR SEQ ID NO:	: 11:	
(:	i) SEÇ	QUENCE CHARACTERIST	TICS:	
	(A) (B)	LENGTH: TYPE:	28 base pairs nucleic acid	

		STRANDEDNESS: TOPOLOGY:	single linear	
(ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 11:	
ATAATCCA	CC TA	TCCCAGTA GGAGAAAT		28
(2) INF	ORMAT	ION FOR SEQ ID NO:	12:	
(i)	SEQ	UENCE CHARACTERIST	ICS:	
	(A) (B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	55 base pairs nucleic acid single linear	
(ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 12:	
AATTTAATI ATGCT	AC GA	CTCACTAT AGGGAGACC	A CACCTTGTCT TATGTCCAGA	50 55
(2) INFO	ORMAT	ION FOR SEQ ID NO:	13:	
(i)	SEQ	UENCE CHARACTERIST	ICS:	
	(A) (B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	30 base pairs nucleic acid single linear	
(ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 13:	
GCACGTAG	rt Ag	CCGGTGCT TATTCTTCA	G	30
(2) INFO	ORMAT	ION FOR SEQ ID NO:	14:	
(i)	SEQ	UENCE CHARACTERIST	ICS:	
	(B)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	53 base pairs nucleic acid single linear	
(ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 14:	
AATTTAATA CGT	AC GA	CTCACTAT AGGGAGAGC	A AGCCTGATCC AGCCATGCCG	50 53
(2) INFO	ORMAT	ION FOR SEQ ID NO:	15:	
(i)	SEQ	UENCE CHARACTERIST	ICS:	



		(A) (B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	32 base pairs nucleic acid single linear	
((ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 15:	
GCTT	rgcgc	CC AT	TGTCCAAA ATTTCCCAC	T GC	32
(2)	INFO	ORMAT	ION FOR SEQ ID NO:	16:	
	(i)	SEQ	UENCE CHARACTERIST	ICS:	
		(B) (C)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	18 base pairs nucleic acid single linear	
((ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 16:	
TCGG	GCCGC	CG AT	ATTGGC		18
(2)	INFO	ORMAT	ION FOR SEQ ID NO:	17:	
	(i)	SEQ	UENCE CHARACTERIST	ICS:	
		(B) (C)	STRANDEDNESS:	40 base pairs nucleic acid single linear	
((ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 17:	
AAC	GCCT:	rt tc'	TTCCCTGA CAAAAGTCC	T TTACAACCCG	40
(2)	INFO	ORMAT	ION FOR SEQ ID NO:	18:	
	(i)	SEQ	UENCE CHARACTERIST	ICS:	
		(A) (B) (C) (D)	LENGTH: TYPE: STRANDEDNESS: TOPOLOGY:	36 base pairs nucleic acid single linear	
((ii)	SEQU	ENCE DESCRIPTION:	SEQ ID NO: 18:	
CGT	AGTTA(GC CG	GTGCTTAT TCTTCAGGT	A CCGTCA	36
(2)	INF	ORMAT	ION FOR SEQ ID NO:	19:	
	(i)	SEQ	UENCE CHARACTERIST	ICS:	
		(A) (B)	LENGTH: TYPE:	46 base pairs nucleic acid	

	(C) STRANDEDNESS: (D) TOPOLOGY:	single linear		
(ii)	SEQUENCE DESCRIPTION: S	SEQ ID NO:	19:	ū
TAATATTA	AC CCTCACTAAA GGGAGACCA	G GCCACTTCC	G CTAACC	46
(2) INF	ORMATION FOR SEQ ID NO:	20:		
(i)	SEQUENCE CHARACTERIST	ICS:		
	(B) TYPE: (C) STRANDEDNESS:	28 base par nucleic act single linear		
(ii)	SEQUENCE DESCRIPTION: S	SEQ ID NO:	20:	
ATAATCCA	CC TATCCCAGTA GGAGAAAT			28
(2) INF	ORMATION FOR SEQ ID NO:	21:		
(i)	SEQUENCE CHARACTERIST	ICS:		
	(A) LENGTH: (B) TYPE: (C) STRANDEDNESS: (D) TOPOLOGY:	55 base par nucleic ac single linear	irs id	
(ii)	SEQUENCE DESCRIPTION: S	SEQ ID NO:	21:	
AATTTAAT ATGCT	AC GACTCACTAT AGGGAGACC	A CACCTTGTC	r tatgtccaga	50 55
(2) INF	ORMATION FOR SEQ ID NO:	22:		
(i)	SEQUENCE CHARACTERIST	ICS:		
	(A) LENGTH: (B) TYPE: (C) STRANDEDNESS: (D) TOPOLOGY:	22 base par nucleic ac single linear	irs id	
(ii)	SEQUENCE DESCRIPTION:	SEQ ID NO:	22:	
GCCGTCAC	CC CACCAACAAG CT			22
(2) INF	ORMATION FOR SEQ ID NO:	23:		
(i)	SEQUENCE CHARACTERIST	ICS:		

8

(A) 20 base pairs nucleic acid single linear LENGTH: (B) TYPE:

(C) STRANDEDNESS: (D) TOPOLOGY:

SEQUENCE DESCRIPTION: SEQ ID NO: 23: (ii)

CCAGGCCACT TCCGCTAACC

20